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NTDB id 123566 FE90 RS00590 WP 002986552.1		MINQWNNLRHKKLKGFTLLEMLLVILVISVLMLLFVPNLSKQKDRVTE	TGNAAVVKLVENQAELYEL	SKG.SKPSLSQ	77				
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS	LVKIHGRKLCQKKQKAFTLIEMLIVLAIIS	TLILLFVPNLIKKKAQVQKT	GEAAVVKVVESQAQLYEL	DHDMDKPNLSE	85			
NTDB id 460 GPW51 RS00790 WP 011921720.1			MKKLVEMKVKGFTLVEMLVVLGIISL	LLFVPNLSQKKDAIQKKGD	AAVVKVVESQMELYEL	EHD.EEATVAD	73		
NTDB id 502 SM12261 RS00955 WP 000738641.1		MKKMMTFLKKA	KVKAFTLVEMLVLLIISV	LLFVPNLTKQKEAV	NDKGKA	AAVVKVVESQAELYSL	DKN.EDASLSK	77	
NTDB id 272 KZH43 RS09255 WP 000738626.1		MKKMMTFLKKA	KVKAFTLVEMLVLLIISV	LLFVPNLTKQKEAV	NDKGKA	AAVVKVVESQAELYSL	EKN.EDASLRK	77	
NTDB id 231 SPD RS09885 WP 000738626.1		MKKMMTFLKKA	KVKAFTLVEMLVLLIISV	LLFVPNLTKQKEAV	NDKGKA	AAVVKVVESQAELYSL	EKN.EDASLRK	77	
NTDB id 197 SPR RS09420 WP 000738626.1		MKKMMTFLKKA	KVKAFTLVEMLVLLIISV	LLFVPNLTKQKEAV	NDKGKA	AAVVKVVESQAELYSL	EKN.EDASLRK	77	
NTDB id 162 SP RS10370 WP 000738627.1		MKKMMTFLKKA	KVKAFTLVEMLVLLIISV	LLFVPNLTKQKEAV	NDKGKA	AAVVKVVESQAELYSL	EKN.EDASLRK	77	
NTDB id 530 SMSK321 RS09230 WP 000738643.1		MKKMMTFLKKA	KVKAFTLVEMLVLLIISV	LLFVPNLTKQKEAV	NDKGKA	AAVVKVVESQAELYSL	DKN.EDASLSK	77	
NTDB id 431 SGO RS09410 WP 012130940.1		MNKLK	KL	RVKAFTLVEMLVLLIISV	LMLLFVPNLTKQKEAV	SDTGNAAVVKV	VESQAELYELKNTG	DQATLSK	74
NTDB id 396 FSA28 RS09395 WP 002263437.1		MKKIK	RVSVRAFTLIEMLVLLIIS	ILLLFVPNLTKQKD	KVKDTGG	SAVVKV	VESQAELYELNKT	.DSATLSK	73
NTDB id 387 SMU RS09025 WP 002263437.1		MKKIK	RVSVRAFTLIEMLVLLIIS	ILLLFVPNLTKQKD	KVKDTGG	SAVVKV	VESQAELYELNKT	.DSATLSK	73
consensus		***	*	***	*****!!!!*!!!!*!*****!*	!!!!*!!!!*!!!!*!!!!	!*	*****	

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NTDB id 123566 FE90 RS00590 WP 002986552.1	LKADGS	ITEKQEKAYQ	DYYDK	HKNEKAR	LSN	108
NTDB id 588 KW2 RS10845 WP 051303741.1	LLS	AGMITQKQVTAYD	DYYDQ	NKNEQ	RNFDD	116
NTDB id 460 GPW51 RS00790 WP 011921720.1	LQAKGY	ITEKQAKQYATAKK				93
NTDB id 502 SM12261 RS00955 WP 000738641.1	LQADGR	ITEEQAKAYKE	YHAKQ	NTSQ	TVAD.	107
NTDB id 272 KZH43 RS09255 WP 000738626.1	LQADGR	ITEEQAKAYKE	YHDK	KNGGAN	RKVND	108
NTDB id 231 SPD RS09885 WP 000738626.1	LQADGR	ITEEQAKAYKE	YHDK	KNGGAN	RKVND	108
NTDB id 197 SPR RS09420 WP 000738626.1	LQADGR	ITEEQAKAYKE	YHDK	KNGGAN	RKVND	108
NTDB id 162 SP RS10370 WP 000738627.1	LQADGR	ITEEQAKAYKE	YND	KNGGAN	RKVND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	LQADGR	ITEEQAKAYN	EYYAK	KNGGAN	RKVND	108
NTDB id 431 SGO RS09410 WP 012130940.1	LVAAGN	ISQKQAD	SYKAYYG	KNNSET	QAVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	LVASGD	ITQEQANSYK	DYYAK	HPSKTR	HRVAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	LVASGD	ITQEQANSYK	DYYAK	HPSKTR	HRVAN	104
consensus	!***!	!***!	***!***	**	** *	

- non conserved
- similar
- ≥ 50% conserved